Serial No.: 09/429,331

### <u>REMARKS</u>

In the Office Communication dated June 13, 2008, the U.S. Patent and Trademark Office (hereinafter "the Patent Office") contends that the application does not comply with the requirements of 37 CFR § 1.821 through 1.825 pertaining to patent applications containing nucleotide sequence and/or amino acid sequence disclosures. In particular, the Patent Office contends that the specification recites sequences in the specification without an assigned SEQ ID No.

In response, applicants respectfully submit that a Response to Sequence Listing Requirement was filed on February 20, 2001, followed by a Supplemental Response to Sequence Listing Requirement on February 27, 2001, in response to an Office Communication of December 5, 2000. Similar to the instant Office Communication, the Office Communication of December 5, 2000 also alleged that the instant application did not comply with the requirements of 37 CFR § 1.821 through 1.825. The responses of February 20, 2001 and February 27, 2001 included amendments to the specification to add SEQ ID NOs. as well as sequence listings, both on paper and in computer readable format. Applicants submit herewith copies of the previously filed responses, as downloaded from the Patent Office Patent Application Information Retrieval (PAIR) system.

As such, applicants respectfully submit that the previously filed responses of February 20, 2001 and February 27, 2001 are believed to address the alleged deficiencies in the specification set forth in the instant Office Communication.

Accordingly, the instant application is believed to be in compliance with 37 CFR § 1.821 through 1.825.

Serial No.: 09/429,331

### **CONCLUSION**

In light of the above remarks, it is respectfully submitted that the present application is now in proper condition for allowance, and an early notice to such effect is earnestly solicited.

If any small matter should remain outstanding after the Patent Examiner has had an opportunity to review the above remarks, the Patent Examiner is respectfully requested to telephone the undersigned patent attorney in order to resolve these matters and avoid the issuance of another Official Action.

### **DEPOSIT ACCOUNT**

The Commissioner is hereby authorized to charge any additional fees associated with the filing of this correspondence to Deposit Account No. <u>23-1665</u>.

Respectfully submitted,

WIGGIN and DANA LLP

Date: 21 July 2008

By:

Todd E. Garabedian, Ph.D. Registration No. 39,197 Attorney for Applicants

10706\86\2072453.1

L

SHERIDAN NEIMARK

ROCER L. BROWDY

ANNE M. KORNBAU

AOI NAWASHIRO

OF COUNSEL IVER P. COOPER JAY M. FINKELSTEIN

NORMAN J. LATKER DIANA MICHELLE SOBO\*

ŀ

BROWDY AND NEIMARK, P.L.L.C.

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TADMITTED IN FL ONLY PRACTICE SUPERVISED BY PRINCIPALS OF THE FIRM

TELEFAX CONTROL SHEET

SENT TO:

laroly

DATE SENT:

SUBJECT:

(including this cover sheet):

FROM:

Remarks:

Attached are the following

1) 2/20/01 Response to Sequence Listing...

and postcord receipt;

2) 2/27/01 Suppl. Resp. to Sequence Listing...

and postcord receipt.

#### CONFIDENTIALITY NOTE

This confidential facelimite message is intended only for the individual entity named above, and may contain information that is privileged and exempt from disclosure under applicable law. If you, the reader of this message, are not the intended recipient, or the employee or egent responsible for delivering this message to the intended recipient, you are hereby notified that you should not copy this facsimile or distribute it to envone other than the intended recipient. In addition, if you have received this telecopy in error, please immediately notify us by telephone or telefax and return the original message to us at the address above via the United States Postal Service. Finally, if it would not inconvenience you, we would appreciate it if you would first refax this message to the intended recipient. Thank you.

If this transmission is not well received, please advise us at our telecopier no. 202-737-3528 or by e-mail at mail@browdyneimark.com, or call our voice telephone no. 202-628-5197.

# IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In Re Application of: PAIGE of at

Application No.: 09/429,331

Examiner: T. Wessendorf

Filed: October 28, 1999

Washington, D.C.

Art Unit: 1627

For: METHOD OF PREDICTING THE ABILITY OF ...

Ally.'s Docket: PAIGE=1D

OR

OR

Date: February 20, 2001

THE COMMISSIONER OF PATENTS AND TRADEMARKS Washington, D.C. 20231

Transmitted herewith is a [ ] Amendment [XX] Response to "Sequence Listing" Requirement with Sequence Listing and Disk and revised pages 239, 244-251, 266-268

in the above-identified application.

- [ ] Small Entity Status: Applicant(e) ctaim small entity status. See 37 C.F.R. §1.27.
- No additional foe is required. ( )
- [XX] The fee has been calculated as shown below:

	(Col. 1)		(Col. 2)	(Cal. 3)			
	CLAIMS REMAINING AFTER AMENDMENT		HIGHEST NO. PREVIOUSLY PAID FOR	PRESENT EXTRA EQUALS			
TOTAL	•	MINUS	20	D			
INDEP.	•	MINUS	з	0			
FIRST PRESENTATION OF MULTIPLE DEP, CLAIM							

_	CIANARE EIGILIA				
		RATE	ADDITIONAL FEE		
]	×	9	\$		
4	×	40	\$		
.]	+	135	S		
ADDITIO	VAL F	FEE TOTAL	s		

SMALL ENTITY

OTHER THAN SMALL ENTITY RATE ADDITIONAL FEE 18 \$ 80 270 \$ TOTAL

- If the entry in Col. 1 is less than the entry in Col. 2, write "0" in Col. 3.
- If the "Highest Number Proviously Pald for" IN THIS SPACE is less than 20, write "20" in this space.
- If the "Highest Number Previously Paid for" IN THIS SPACE is less than 3, write "3" in this space.

The "Highest Number Previously Paid For" (total or independent) is the highest number found from the equivalent box in Col. 1 of a prior amendment of the number of claims originally filed.

Conditional Pathlon for Extension of Time

If any extension of time for a response is required, applicant requests that this be considered a poblion therefor.

[XX] It is hereby politioned for an extension of time in accordance with 97 CFR 1.136(a). The appropriate fee required by 37 CFR 1.17 is calculated as shown below:

\$1.16 and all palent processing fees under 37 CFR §1.17 (broughout the prosecution of the case. This blanket authorization does not include patent issue fees under 37 CFR §1.18.

	Small Entity	Other Than Small Entity
	Responso Filed Within	Response Filed Within
	( ) First - \$ 55.00	[ ] First - \$ 110,00
	[ ] Second - \$ 195,00	[XX] Second - \$ 390,00
	[ ] Third - \$ 445.00	[ ] Third - \$ 890,00
	[ ] Fourth - \$ 695.00	[   Fourth - \$ 1390,00
	Month After Time Period Set	Month After Time Period Set
[ ]	[ ] Less (ees (\$) already paid formonth(s) extension of time.  Please charge my Deposit Account No. 02-4035 in the amount of \$	•
pox	Credit Cerd Payment Form, PTO-2038, is attached, authorizing payment in the	on same of \$390.00.
[ ]	A check in the amount of \$ is attached (check no. ).	
bxd	any Extension of Time fee, not covered by check or specific authorization, but	nal foes which may be required in connection with this application or credit any tile not limited to payment of all fees associated with this communication, including is also intended to include all fees for the presentation of extra claims under 37 CFR ecution of the case. This blanket authorization does not include astertions.

BROWDY AND NEIMARK

Altomeys for Applicant(s)

er P. Coope Registration No. 28,005

(202) 737-3528 (202) 628-6107

Fecsimile:

(NOVA) APPLICANT(S): PAIGE of all APPLICATION NO: 09/429,331	DOCKET NO.: PAIGE = 13
THE PATENT AND TRADEMARK OFFICE STAMP HEREON ACKNOWLEDGES RECEIPT OF THE FOLLOWING PAPERS:  FEES \$ 340.00	SECTIVED
☑ PTO FORM 203B ☐ (CH. #	
☐ MISSING PARTS RESPONSE WITH DECL ☐ AMENDMENT ☐ PRELIMINARY ☐ SUPPLEMENTAL ☐ REPLY TO OFFICE ACTION ☐ RESTRICTION/ELECTION REPLY	ASSIGNMENT INFORMATION DISCLOSURE STATEMENT FORM 1449 & PATENTS/PUB: PRIORITY DOCUMENT(S) NO
SEQUENCE LISTING MOTH DISK REFORM TRANSMITTAL (clicle one) NOTICE OF APPEAL APPEAL BRIEF (TRIPLICATE)	☐ DECLARATION UNDER § ☐ LETTER TO DRAFTSMAN ☐ SHEETS OF DRAWINGS ☐ ISSUE FEE TRANSMITTAL FORM ☐ MAINTENANCE FEE LETTER
PREPLY BRIEF (TRIPLICATE)  TOTHER SELVINGE CASES 239, 24  Copy of Notice to Comply  Ban-2	+- 251, 266-268, 270 and 272;

From-BROWDY NEIMARK

Application No.: (

T-107 P.008

# NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

wing reason(s):	
1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.	
2. This application does not contain, as a separate part of the disclosure on paper copy, a "S quence Listing" as required by 37 C.F.R. 1.821(c).	
3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).	•
4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked-up "Raw Sequence Listing."	
5. The computer readable form that has been filed with this application has been found to be damage and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).	∌d
6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).	
7. Other:	
LI .	
Applicant Must Provide:	
Applicant must read able form (CRF) copy of the "Sequence Listing".	
An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.	
A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).	
For questions regarding compliance to these requirements, please contact	
For Rules Interpretation, call (703) 308-4216 For CRF Submission Help, call (703) 308-4212 For Paleotin software help, call (703) 308-6856	
TOT THE METER AND THE PERPONSE	

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE

# IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:

PAIGE et al

Appln. No.: 09/429,331

Filed: October 28, 1999

For: METHOD OF PREDICTING THE ABILITY OF COMPOUNDS TO MODULATE THE BIOLOGICAL ACTIVITY OF RECEPTORS

ART UNIT: 1627

Examiner: T. Wessendorf

Washington, D.C.

February 20, 2001

Atty.Docket: PAIGE=1D

RESPONSE TO "SEQUENCE LISTING" REQUIREMENT

Honorable Commissioner of Patents Washington, D.C. 20231

sir:

In response to the Notice to Comply, mailed December 5, 2000, petition for a two-month extension of time and payment of late fees attached hereto, please amend the application as follows:

## IN THE SPECIFICATION

-- (SEQ ID NO:4) --.

Page 132, line 14, before "biotin" insert

--(SEQ ID NO:1)--.

Page 136, line 26, after "...CTGCG" insert

--(SEQ ID NO:3)--;

line 27, after "...ACCTA" insert

In re Appln. No 09/429,331

Page 150, line 22, after "...GTCAG" insert -- (SEQ ID NO:5) --;

line 25, after "...GTCAG" insert

-- (SEQ ID NO: 6) --;

line 28, after "...GTCAG" insert

-- (SEQ ID NO:7)--;

line 31, after "...GTCAG" insert

-- (SEQ ID NO:8) --;

line 33, after "...TCGAG" insert

-- (SEQ ID NO:9) --.

Page 162, line 33, after "...CAGT-3'" insert

-- (SEQ ID NO:14) --;

line 36, after "...TAGA-3'" insert

-- (SEQ ID NO:15) --.

Page 173, line 26, after "...SLLSR" insert

-- (SEQ ID NO:187)--.

Page 183, line 6, after "SRLXXLL" insert

-- (SEQ ID NO:2) --.

Page 225, line 4, after "...KQAV" insert

-- (SEQ ID NO:10) --;

line 5, after "...GVSR" insert

-- (SEQ ID NO:11) --;

line 6, after "...MLSR" insert

-- (SEQ ID NO:12) --;

line 7, arter "...YASR" insert

May-15-2002 11:56

-- (SEQ ID NO:13) --.

Page 238, line 2, after "...GHSR" insert

-- (SEQ ID NO:59) --;

line 3, after "...WRSR" insert

-- (SEQ ID NO: 60) --;

line 4, after "...KDSR" insert

-- (SEQ ID NO: 61) --.

Attached are copies of pages 239, 244-251, 266-268, 270, and 272 in which sequence identifiers are marked in red. Entry of these revisions is respectfully requested.

Please enter the enclosed "Sequence Listing", pages 1-79.

### REMARKS

- Applicants hereby submit the following: 1.
- [XX] a paper copy of a "Sequence Listing", complying with \$1.821(c), to be incorporated into the specification as directed above;
- ] an amendment to the paper copy of the "Sequence Listing" submitted on , the amendment being in the form of substitute sheets;

In re Appln. No-09/429,331

- [XX] the Sequence Listing in computer readable form, complying with \$1.821(e) and \$1.824, including, if an amendment to the paper copy is submitted, all previously submitted data with the amendment incorporated therein;
- [ ] pursuant to \$1.821(e), reference is made to the computer readable form filed on , in USSN , which presents the identical Sequence information, the use of which is now requested, in lieu of submitting a new computer readable form; and/or
- [ ] a substitute computer readable form to replace one found to be damaged or unreadable.
- [XX] 2. The description has been amended to comply with \$1.821(d).
- 3. The undersigned attorney or agent hereby states as follows:
  - (a) this submission is not believed to include new matter [\$1.821(g)];

- (b) the contents of the paper copy (as amended, if applicable) and the computer readable form of the Sequence Listing, are believed to be the same [\$1.821(f) and \$1.825(b)];
- (c) if the paper copy has been amended, the amendment is believed to be supported by the specification and is not believed to include new matter [\$1.825(a)]; and
- (d) if the computer readable form submitted herewith is a substitute for a form found upon receipt by the PTO to be damaged or unreadable, that the substitute data is believed to be identical to that originally filed [\$1.825(d)].

Respectfully submitted,

BROWDY AND NEIMARK

Attorneys for Applicant(s)

By:

Tver P. Cooper

Registration No. 28,005

IPC:al
624 Ninth Street, N.W.
Washington, D.C. 20001
Telephone No.: (202) 628-5197
Facsimile No.: (202) 737-3528
F:\,N\Nova\PaigeID\Pto\SequenceResponse.doc

239

Table 3: Phage/Peptide Classification # and isolation mathod E? + estradiol ER + estradiol #4 <u>class l</u> SSNHQSSRLIELLSR 62 SRLKELLLLPTDLSR 63 En + estradiol #15 SSKLYCLLDESYCSR W ER + estradiol #35 #41 ER + estradiol 65 HGPLTLNLLRSSGG SRLEYWLKWEPGPSR 66 **#12** 

class 2

SSCKWYEKCSGLWSR 67 #7 Ξ3 ER + estradiol SSEYCFYWDSAHCSR 65 #33 **Ξ**R #31 #24 ER + estradiol 69 SSWVLLRDLPWGSR SSWVRLSDFPWGVSR 70 10

ER + estradiol Class 3 SSLTSRDFGSWYASR 71 #5

Class 4 15

ER SRTWESPLGTWEWSR 72 #13

Class 5

#48 ER SAACATISHYLMGG 73

. 22

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1.5

	present when present when peptide was identified Tamoxifen
hh7.	Serve in the pepting presence of pepting presence of pepting receptor form in the pepting receptor form in the pepting receptor form in the pepting receptor form receptor form receptor form receptor form receptor recept
	Peptide Sequence Peptide Sequence Sequence Sequence Sequence Sequence Sequence Sequence Service Servic
	<u>rable 7: New Era</u> Peptide  name  1PT  2PT  4PT  4PT  7PT  10PT  11PT  12PT  13PT  14PT  15PT  16PT  18PT

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Estradiol Estradiol Estradiol Estradiol Estradiol **Estradio**l **Estradio**] Estradiol Estradiol Raloxifen Raloxifen Tamoxifen Buffer Buffer Buffer Tamoxifen Tamoxifen Buffer Buffer Tamoxifen Buffer Estradio1 Buffer Buffer Estradiol Estradiol Estradiol present when peptide was identified H SERM Table 8: New ER $\alpha$ -ERE Peptide Sequence Information מססט ರ ed in the No. presence of ਨ ℧ receptor 8 ਠ ช 8 form SRICFGDWCWLGGVDVLSR 109 SSINMVVDTPWCGKWVCSR 110 SSDWCFGWGGWCASEAVSR II7 SRNWDWAALELLPYPHPSR 11 SRVFGVSGGEVVLINGSSR 10% SRLEELLLMDFWRSR (25 SKLPPSVFSMCGSEVCLSR 106 SSKLWQLLSSPIDSR 126 SSICLYCLLDESYCSR [2] 当主 SRLTCLIASNGWDSEQCSR 103 SSLTSRDFGSWYASR 104 HSHNHHSPWLFRLLGG 100 SRFEIWKPEPGCVSSLENWE peptide sequence 300 SRLVALLKSPWSVSR SSLITSRDFGSWYASR SSTGILMKLLTAESR SSHGILWRLLSEGSR SRSDSILWRMLSESR SSRPDAAFFGAKLSR SRSPILTHLLSLGSR SRSYHGEWGVWTLSR SSRIADLFWRLEPSR SRWNDTSWWLEELSR HSHPLPPLLSRLLTGG SSRPSESFWEKQLSR SSRPTAEWFRENLSR HSHPHHSHLLYKLMGG SRILQLDWGTLYSR PGKRVCSR 6E 11.12 10B T1-10 T2-10 T3-11 14-10 **6B** 9B **2B** 1B 7.8 2R 3B E1-7 T3-1 43 E1-4 Pepti H E1-3 E1-1 de name

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S

Estradiol Estradiol Estradiol Estradiol Estradiol Estradiol

SRLLMDMLMSDDYVTVSR 128
SSRLLACELMYEDADVCSR. 129
HSHSPLLMALLAPPGG 130
SRLEYYLRLGTYESR 131
SSCLREILLYGACSR 131
SSRTAEDYCFFADDYWCSR 131
SSLRCYLSSKYDQWACSR 131
SSLRCYLSSKYDQWACSR 131
SSYKPHSLLEWHLLGGTSR 135

From-BROWDY NEIMARK

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Buffer Buffer Buffer Buffer Buffer Busser Busser Buffer Buffer Buffer Buffer Buffer Buffer Buffer Buffer peptide was present when identified Table 9: New ERB-ERE Peptide Sequence Information Poplide Sequence Seq. of receptor loral lead SSEACYGRWMLCEQLGVSR 145 HSWLOPWRLSSIDLGG 157 3 3 <u>8</u> 165 SSQVWPGPWRLVESR 146 SRGGGECLOPWCLSR 144 SSTSWLIHYLMGTSR 143 SSGPWLMHYLGGGSR 142 SSGLPPNFERMLKSR 147 SSWPNPTFWERQLSR 138 HSYSSHPLLLSYLWGG SSWPGAEWFKEQLSR SSKLYCLLDESYCSR SSSLGRWRLSELESR SRECVGGWCLAELSR SSIPPRSWWLSQLSR SSSOPWRWGLSIESR SSSMMREPFERELSR SRLHCLLDSSYCSSR SRLHCLLDSSYCSSR SYSKEWFEERLNSR 2413-13 23B-B 21B-D 20Љ-В 17B-B 18B-B 193-6 16B-ß 14B-B 12B-9 8B-B 9B-B 7B-B 69-69 SB-B 4B-β 3В-6 2B-β 1B-B

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S

Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Estradiol Tamoxifen Tamoxifen .Famoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Busser Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen Tamoxifen 248 SKI)LYSQMINEFFQINI,SR 175 SSRLVPHSFWLDGLMHGSR 173 SSISTYHMGEWFYAMLSSR 17# SRPGCLRGY.WCLADTPIPSR 172 SRLNGVFCHDSSDLWYCSR [7] SSGPFYVGGMLWPADCLSR 169 SSDLINDCLGVWCLSR 170 SRMFQVCGDEVCLRSR 169 Sregwmgpwrladsr 165 SSWASREWWVRELSR 168 SRLPPS.VFSMCGSEVCLSR (6) SSPGSREWFKDMLSR 167 SSQEEWLLPWRLASR 162 SSARPWWLQFEGSSR 161 SREWEDGEGGRWLSR 158 SSRGLLWDLLTKDSR SRNECIGPWCLTISR SRYCLGDVWCLDSR HSTDMGWLRPWRLLGG SSTITMEDFFYERLSR SSWINSREFFLSQLSR SSVFTIMDGKVALSR 23.1.-13 22T-β 20T-B 21T-B 18T-B 17T-β 15T-B 16T-B 14T-B 12T-β 13T-β 10T-B 9T-B 8T-B  $7T-\theta$ 6T-β 4T-B 5T-β 2T-β IT-β

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Estradiol
6 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5
SRHGILWDLLGGDSR 178 SRLHDLLLRDESPSR 178 SRDWRSGFLYELLSR 179 SSDTRSRLYELLSSSYTSR 180 SSTEELLRVGVLTSR 181 SRLEELLRVGVLTSR 187 SSTEELLROGGSAETV 187 SSILERLLGGGSAETV 187 SSIKDFPNLISLLSR 186 SSGSSAGRLMMLLQDGSR 186 SSGSSAGRLMMLLQDGSR 187 SSHCLLLDAGGCSR 191 SSRLLCLLDAGGCSR 191 SSRLLCLLDAGGCSR 192 SSLKCLLOSSPQKQPFCSR 192 SSLKCLLOSSPQKQPFCSR 196 SSAGLLEMNLGGSR 196 SSAGLLEMNLSRSR 196
2E-B 3E-B 4E-B 6E-B 6E-B 7E-B 10E-B 11E-B 13E-B 14E-B 15E-B 17E-B 17E-B 20E-P 20E-P 20E-P 21E-B

Estradiol Estradiol Estradiol Estradiol Estradio Estradio Estradiol Estradio Estradiol Estradiol Estradio Estradiol SSRPSFTIPWWFDDPSRSR 208 SSLDLSQFPMTASFLRESR 210 SSTRLMCWLGSADTSHCSR 203 SSYDWQCPSWYCPAPPSSR 204 SSYEIPKWALQWLSR 209 SSRTLYCHLTSSNPEWCSR 202 SSQWQAPWWYIDASR 201 SSWDFRVPWWYNNSR 206 SSTTWRCPEWYCGSR 205

SRWWLDDHELLLYSSR SSRTLLEHYLLGGSR SRPEGSSWLLHYLSR

250.

34E-B 35E-B 33E-B 27E-B 28E-B 29E-B 30E-B 32E-B 31E-B 25E-β 26E-B

Table 10: Panel Peptides for Example 2

Alternative name paronthesized. Modulator used to isolate peptide in brackets. BIII, SSEACVGRWMLCEQLGVSR. (B3) [no modulator] (Set 12 10 1721) α/β III, SSWDMHQFFWEGVSR (AB3) [4-OH lamoxifen] (SEB 10 NO: 213) alp IV, SRLPPSVFSMCGSEVCLSR (ABA) [same] (SEG ID MOI 214) B 11, SSLDLSQFPMTASFLRESK (B2) [1714-extradiol] (Seg 15 rues 220) (SER 10 PM: 211) B 1, SREWEDGFGGRWISR (B1) [4-Ohl tamoxifen] (Sea to not 219) (32 mm or 681) α III, SRTWESPLGTWEWSR (A3) [no modulator] (Less 12 por: 21 8) a II, SSLTSRDFGSWYASR (A2) [17β-estradiol] (1602 10 pol: 217) α 1, SSEYCFY WDSAHCSR (A1) [17fp-estradiol] (Seig 13 two: 216) (SER 18 MG: 212) a/B 11, SAPRATISHYLMGG (AB2) [no modulator] a/b 1, SSNHQSSRLIELLSR (AB1) [17f3-cstradiol] α/β V, SSPGSREWFKDMLSR (ABS) {same}

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ser is no:
         SRAGLĻSDLLEGKSR
 Table 100
                              223
         SSRSLLRDLLMVDSR
 A
         SSNKLLYNLLKMESR
                               224
          SSNKLLLNLLSTPSR
SSKSLLLNLLSTPSR
     HSFPRESLLVRLLQGG
              SRLEMLLRSETDFSR227
5.
               RLEELLKWGSVTSR 228
               RLEQLLKEEFSYSR229
               R'LEQLLRSEPDFSR230
               RLEDLLRAPFTTSR231
              S
               RLESLLRFGQLDSR 232
10
              S
               RLLSLLVGDFNSR
              s
               SRLEELLLGTNRDSR23Y
             S.S
               SRLEELLLMDFWRSR 235
                RLKELLLPTDL
                                 S R 236
                                 S R 237
               SRLECLLEGRLNC
15
               S
             SSKLYCLLDESYCSR 238
                RLSCLLMGFEDCSR239
               SKLIRLLTSDE E.LSR 240
               SRLMELLQEGQGWSR 241
20
               SRLIELLSR 242
               SRLWQLLASTDTSR 26
       SSNHQS
               SRLWQLLSSPIDSR 245
SKLWQLLSSPIDSR 245
              S
                SRLVĀLLKSPWSVSR
              S
             SNSMLWKLLAAPSR 246
             SSKTLWRLLEGERSR 247
 25
             AGPVLWGLLSESR 248
             RSPILTHLLSLGSR 249
             STGILWKLLTAESR 250
SHGILWRLLSEGSR 251
 30
                  KLVQLLTTTAE 251
                  ILHRLLQEGSP 253
    В
                   LRYLLDKDEK 254
                  LLQQLLLTE 255
                  L
        SRCla
  35
                  QLSELLRGGSG 256
QLVLLLHAHKC 257
         CBP
                   YLEGLLMHQAA 258
                    LASLLQSESS259
                    LKTLLKKSKV260
                    LALLISSEAH 261
   40
                   Н
                    LLHLLKSQTI 262
                   Q
         RIP140
                   LLQLLLGHKNE 243
                   A T O'T T T G M B K G 57%
                   LLS.RLLRQNQD 265
                   VIKQLLLSENC 266
   45
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SRCla = human steroid receptor coactivator la,

CBP = mouse cAMP-responsive element (CREB)-binding

protien,

RIP 140 = human RIP140

May-15-2002 12:02

Table 101	SEO IBNO:
5	Class I ER4 SSNHQSRLIELLSR 267 D2 GSEPKSRLLELLSAPVTDV 280 D30 HPTHSSRLWELLMEATPTM 281 D11 VESGSSRLMQLLMANDLLT 282
	Class II
10	D47 HVYQHPLLLSLLSSEHESG 269 C33 HVEMHPLLMGLLMESQWGA 269 D14 QEAHGPLLWNLLSRSDTDW 270
<b>-</b>	Class III F6 GHEPLTLLERLLMDDKQAV 271 D22 LPYEGSLLLKLLRAPVEEV 272 D48 SGWENSILYSLLSDRVSLD 273 D48 AHGESSLLAWLLSGEYSSA 274 D43 AHGESSLLAWLLSGEYSSA 275
15 .	D17 GVFCDSILCQDLLAGSDAPS 276 D41 HHNGHSILYGLLAGSDAPS 277 D26 LGERASLLDMLLRQENPAW 237 D40 SGWNESTLYRLLQADAFDV 218 D40 SGWNESTLYRLLQADAFDV 219
. 20	D15 PSGGSSVLEIDHIM F4 PVGEPGLLWRLLSAPVERE 281
	Misc. D10 WEEHSQMLLHLLDTGEAVW6 283
Erßsp	#293 SSIKDFPNLISLLSR [87
25 GRIP-	NR1 DSKGQTKLLQLLTTKSDQM 10 NR2 LKEKHKILHQLLQDSSSPV 17 NR3 KKKENALLRYLLDKDDTKD 18
src-	NRI YSQTSHKLVKLLTTTAEQQ 19 NR2 LTARHKILHRLLQEGSPSD 20 NR3 ESKDHQLLRYLLDKDEKDL 21
30 .	

		· (T-P	eptide	:5)
	•	Sequence/Motif Aligned SEALLITHSEFLDSHTK 22 1	L	ibrary
	anan Gial GD	GIFYS Aligned AFE	<u> </u>	E
	Table 202A.	SRAHLLTWSEFLDSHTK SRAHLLTWSEFLDSHTK	BUF	=
	<u>ID</u>	SRAHLLINGEL GOLNP 23	Buf	
	99	SSGELITWYEFLGDLNP 23	BÜF	, <u>E</u>
	103	COCET THE LIBRAIC	GTP	<u>K</u>
		PET TWMEL TOD	GTP	<u> </u>
5	107	ALMADE TWED IN CO.		CWL R Y PHD12
	361	TO STATE OF THE ST	BUF	v
	388,391	contradot al Région	GTP ·	2013
	45	NLMIWIE I III 29 F	BUF	PHUIZ
	297.401,412	* PDI WINGER DI	GTP.	<u>N</u>
_	15r2,301,394	VTVSLYEFTED	BUF	. <u>Н</u>
JO	380,381,140	OCCULATIONE SI	,,,,,	
	16	SSRGEYWWEL EGGE 33	BUF	
	360	SSAUGIFWWIII		
	101	- 24	GTP	
		LGRGTTDMPPWAWWS	GDP	
	275, 123, 125, 247	CHOMED DWOWIN -	BUF	
15	272 234	" TO ALD DIKMY'I. JA	_	
		TARACCIME VIII O		
		THE MOWEMRVVE 38	GIR	
	387	FEGUTIAL CO.		
	386			
15	101 375,123,125,247 331,334 37 387	LGRGTTDMPPWAWWS 3 7  NYTERPWVWYH 35  SSLYSMEPWKWYT 36  KWWESDWFVNFG 37  EEGMDWFMRVVE38	_	

Mar-15-2002 12:02

		r crp-specific Phage			
	Table 202B: Glo	1 GTP-Specific Phage	60 ID 1	<b>10:</b>	
	(T-Peptides)	• -			
	370,377,378 244	SVLSSEMCFGWACY SEMCFGWACY FNEVCLGWQCY	39 40 41	GTP	M · <u>PARO</u> <u>K</u>
5	366,G12	SSNARPCQGWHCYL	SOSR	42	
>	G33, G34	WDGGVWMGPAS Y		GTP	<u>K</u>
	353	MGGGVWRGP 4	Į.	GTP	<u>Y</u>
	408	GDACCAMPG SEGNO	R <b>45</b>		
	G22,G25	SSWDGGVWWGQYGSR	. <b>46</b>		
10	G11,G26-29 G9,G10	SSNLDGCFTSGGVWSGCSR LGYDINGVWIG	47 48	GTP	N
	382	ICDIIPWEESCSR	49	GTP	<u>ė</u>
	384	ACGPAICPWDFMPQL	50	GTP	PARO
	413				

Note: clone 244, which was identified in a screen for peptide which bound GDP:G-alpha, is suspected to having increased the affinity of the G-alpha for GTP through a conformational change.

T-097 P.022/027 F-749

	m-bas anace Girl	GDP-Specific Phage				
	(D-Peptides)		SED IDAR	-		
		SRGPQLTWQEFLTGAASSR	গ্	<b>~</b>		
	G4	nvvtyweflgp	<u>52</u>	GD?	1	•
	314	SREFVTWKEFLGS	<i>5</i> 3	BUF	K	
5	73	SOLTWREFLFG	54	Gシラ	R	
	343	SSHLMTWHEFISD	55	<b>GDP</b>	H	
	217	. SRDGFETWAEFLGASGS	56	BUF	_	
	93	SRLTWSEYLSEIDP	51	BUF	CML	•
	62	SRTVTWVDFLKET	<i>5</i> 8	GDP	D	
10	193	MSWYZFMTEESM	SBS	GDP	CWI	
	324	AKHDLSWYEFLQLPI	286	GTP	V	
	400	SRLSWWEFLGASDCGTC	287	GDP X	14C <w></w>	
	281	DLLSLKEFLAT	288	GTP	K	
	359,161	SSPNLLTLEEFLS	287	GDP	L	
15	176	KTYSLYEFLEL	290	GTP	N	
	380,381,140	MSNRYTIYEFLNLHS	291	GTP	Y	
	409,24x2	LHWWEVLAEK	292-	GDP	CWL	
	320	SSPQPLLHWWENMTEPP	293	GD₽	<ul> <li>KNK</li> </ul>	
	230	SRAGESVHWWEVL	294	GDP	H	
20	213	RAGESVHWMEYIATL	29.5	GDP	N	
	266	EMISWHQYLLSI			GDP	PARO
	237		7A 287	BUF	М	
	126,128,133,242	VPWWVWLAEGD	248	GTP	N	
	379	SREIYWWDWLTDT	299	GD₽	. D	
25	196	FGSNMLDLPTFLDWL	300			.0
	117	FGSNMLDHFIFHDMA SRITFWELMLEGG	301			
	92	SRTPYEWLGYWGA	302			
	179	SKIFIEMESIKOM		-		
		YDMCTWLEFLDGGEC	262	GDE	X14	LCM
	289	SPLCTWAEYLMEPSC	304	GDE	N	
30	265	TOWCTWAEFLSSTDC	305			•
	273	SSDGCTWQEFLAGHGPC			P N	
	272,282,6R2	22Ddc14/5=1 7:101101 -		-		
	•	PFNNPPWMWWS	30	7 GD:	P P	
	337,339	SSPTVHENI PPWLWWSP	30	g GD	N g	
	268	CIHADAMAMQE	30		P P	
3	5 330	. GEDABBAAMDE	3 r		P P	
	329	YSQVFGDAPVWAWYSSR	31	I GD	P XI	4CW
	280	YSQVFGDAPVWAW155K WTPSDWQWWRSK	31	کہ GE	DP CW	
	319	WIPSDWQWWSX	31			ARO
	115	SSHWSSDSIFPGFWYSG	-			
		SRGGVDLDIGNSA	31	ur G!	Q QC	
4	10 197	EGEDVRTRIAN	3		DP R	
	347	EGEDAKIKIAN				

### SEQUENCE LISTING

<110> PAIGE, Lisa A. MCDONNELL, Donald P. CHANG, Ching Yu NORRIS, John HAMILTON, Paul T. FOWLKES, Dana M. BARNETT, Tom CHRISTIANSEN, Dale J. BUEHRER, Benjamin

- <120> METHOD OF PREDICTING THE ABILITY OF COMPOUNDS TO MODULATE THE BIOLOGICAL ACTIVITY OF RECEPTORS
- <130> PAIGE1D
- <140> 09/429,331
- <141> 1999-10-28
- <150> PCT/US99/06664
- <151> 1999-03-26
- <150> 60/082,756
- <151> 1998-04-23
- <150> 60/099,656
- <151> 1998-09-09
- <150> 60/115,345
- <151> 1999-01-08
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Gly ser Gly Lys

<210> 2

<211> 8

<212> PRT

<213> Artificial Sequence

Mat-15-2002 12:03

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<210> 3 <211> 23 <212> DNA <213> Xenopus laevis	
<400> 3 gatctaggtc acagtgacct gcg	23
<210> 4 <211> 23 <212> DNA <213> Xenopus laevis	
<400> 4 gatccgcagg_tcactgtgac cta	23
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<210> 6 <211> 73 <212> DNA <213> Artificial Sequence	
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<220>
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gaetgtgcga attetettet tgggatatgc atcaattttt ttgggaaggt gtttctagac 60
togagogtgt cag
<210> 8
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tegagegtgt cag
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                                       10
 Gln Ala Val
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                                     10
                  5
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  <223> N at each occurrence is A, C, G or T; K at each
        occurrence is C or T
  agtgtgtgcc tegagannkn nknnknnknn knnknnkctg nnknnkctgc tgnnknnknn 60
  knnknnknnk nnktctagac tgtgcagt
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Ala Glu Gln Gln

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     3' end of SEQ ID NO:14
                                                                  .15
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actgcacagt ctaga
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 Asp Ser Lys Gly Gln Thr Lys Leu Leu Gln Leu Leu Thr Thr Lys Ser
 Asp Gln Met
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 Leu Lys Glu Lys His Lys Ile Leu His Gln Leu Leu Gln Asp Ser Ser
  Ser Pro Val
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   Thr Lys Asp
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       _ 5
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 Lys Thr Tyr Ser Leu Tyr Glu Phe Leu Glu Leu
                                       10
                  • 5
   1
  <210> 31
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  <213> Artificial Sequence
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                                        10
                    5
  <210> 32
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<211> 15
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 Ser Ser Arg Gly Glu Tyr Trp Trp Glu Phe Leu Gly Tyr Ser Arg
                                       20
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Ser Ser Ala Asp Gly Ile Phe Trp Trp Glu Tyr Ala Arg Glu Ala Gly
                                      10
Glu
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Leu Gly Arg Gly Thr Thr Asp Met Pro Pro Trp Ala Trp Trp Ser
                  5
                                      10
                                                       15
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<400> 35
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Mar-15-2002 13:09 From-BROWDY NEIMARK
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<211> 13
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              . 5
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<211> 15 <212> PRT

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Gln Ser Arg
 <210> 43
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  1
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<220>

From-BROWDY NEIMARK

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Met Gly Asp Ser Val Leu Pro Tyr Gly Gly Val Trp Leu Gly Pro
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Ser Ser Asn Leu Asp Gly Cys Phe Thr Ser Gly Gly Val Trp Ser Gly
Cys Ser Arg
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Leu Gly Tyr Asp Ile Asn Gly Val Trp Ile Gly
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Ser Ser Arg
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                                      10
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. Ser
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  <211> 15
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Ser Ser Val Thr Lys Lys Ala Leu Thr Ile Ala Lys Asp Ser Arg
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                   5
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<211> 15

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Ser Ser Cys Lys Trp Tyr Glu Lys Cys Ser Gly Leu Trp Ser Arg
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<210> 78
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      peptide.
<400> 78
Ser Ser Arg Met Gly His Val Trp Tyr Asp Trp Thr Phe Ser Arg
<210> 79
<211> 15
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<400> 79
Ser Ser Arg Leu Leu Gly Asp Phe Gly Gly Ser Val Val Ser Arg
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<212> PRT
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Ser Ser Lys Tyr Val Phe Gly Phe Gln Val Ala Gly Gly Ser Arg
                                      10
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 <210> 81
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Ser Ser Trp Ala Gly Ile Lys Phe Gly Lys Pro Pro His Ser Arg
<210> 82
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Ser Arg Asp Thr Gly Asp Met Trp Trp Gly Arg Gly Gly Ser Arg
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Ser Ser Gly Arg Tyr Asp Pro Phe Val Leu Asn Ala Ala Ser Arg
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<220>

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Ser Ser Trp Pro Tyr Leu Pro Lys Arg Glu Glu Trp Ala Ser Arg
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ser Ser Gly Trp Ile Glu Gln Lys Leu Arg Gly Ser Phe Ser Arg
<210> 88
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<210> 89
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<400> 93 -

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Ser Ser Ser Thr Pro Gly Trp Trp Glu Trp Asp Trp Ala Ser Arg
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<210> 94
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Ser Ser Tyr Trp Asp Gly Ser Trp Arg Arg Lys Glu Thr Cys Val Ser
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Cys Ser Arg
<210> 95
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<212> PRT
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Ser Ser Arg Thr Ala Glu Asp Tyr Cys Phe Phe Ala Asp Asp Tyr Trp
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                                      10
Cys Ser Arg
<210> 96
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       peptide
 <400> 96
 Ser Ser Arg Ala Leu Ala Leu Phe Pro Val Gly Met Glu Ser Arg
                  5
 <210> 97
 <211> 19
 <212> PRT
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Ser Ser Asp Cys Glu Ser Leu Thr Ser Tyr Pro His Leu Lys Ala Leu
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Cvs Ser Arg
<210> 98
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<400> 98
Ser Ser Thr Ala Thr Ala Leu Arg Asp Arg Leu Ala Tyr Ser Arg
                                      10
<210> 99
<211> 15
<212> PRT
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 Ser Ser Gly Lys Thr Arg Glu His Tyr Arg Glu Gly Thr Ser Arg
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 <210> 100
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 <400> 100
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 <210> 101
 <211> 16
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<213> Artificial Sequence
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      peptide
His Ser His Pro His His Ser His Leu Leu Tyr Lys Leu Met Gly Gly
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      peptide
 <400> 102
 His Ser His Pro Leu Pro Pro Leu Leu Ser Arg Leu Leu Thr Gly Gly
                                       10
                   5 .
 <210> 103
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 Ser Arg Leu Thr Cys Leu Leu Gln Ser Asn Gly Trp Asp Ser Glu Gln
 Cys Ser Arg
  <210> 104
  <211> 15
  <212> PRT
  <213> Artificial Sequence
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  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 104
  Ser Ser Leu Thr Ser Arg Asp Phe Gly Ser Trp Tyr Ala Ser Arg
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  <210> 105
<211> 14
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 105
Ser Arg Thr Leu Gln Leu Asp Trp Gly Thr Leu Tyr Ser Arg
<210> 106
<211> 19
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Leu Pro Pro Ser Val Phe Ser Met Cys Gly Ser Glu Val Cys
                                      10
Leu Ser Arg
<210> 107
<211> 28
<212> PRT
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      peptide
Ser Arg Phe Glu Ile Trp Lys Pro Glu Pro Gly Cys Val Ser Ser Leu
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Glu Asn Trp Glu Pro Gly Lys Arg Val Cys Ser Arg
                                  25
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<210> 108
<211> 19
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
    .peptide
Ser Arg Val Phe Gly Val Ser Gly Gly Glu Val Val Leu Ile Asn Gly
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<211> 15 <212> PRT

<213> Artificial Sequence

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Ser Ser Arg
<210> 109
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <220>
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 Ser Arg Leu Cys Phe Gly Asp Trp Cys Met Leu Gly Gly Val Asp Val
                                       10
 Leu Ser Arg
 <210> 110
 <211> 19
 <212> PRT
<213> Artificial Sequence
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 <223> Description of Artificial Sequence: Arbitrary
      peptide
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 Ser Ser Leu Asn Met Val Val Asp Thr Pro Trp Cys Gly Lys Trp Val
 Cys Ser Arg
 <210> 111
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
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 <400> 111
 Ser Ser Arg Pro Asp Ala Ala Phe Phe Gly Ala Lys Leu Ser Arg
 <210> 112
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<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 112
Ser Ser Arg Pro Ser Pro Ser Phe Trp Glu Lys Gln Leu Ser Arg
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Ser Ser Arg Pro Thr Ala Glu Trp Phe Arg Glu Asn Leu Ser Arg
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Ser Arg Trp Trp Asp Thr Ser Trp Trp Leu Glu Glu Leu Ser Arg
<210> 115
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 115
Ser Ser Arg Ile Ala Asp Leu Phe Trp Arg Leu Glu Pro Ser Arg
<210> 116
<211> 15
<212> PRT
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 <223> Description of Artificial Sequence: Arbitrary
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peptide

<400> 116 Ser Arg Ser Tyr His Gly Glu Trp Gly Val Trp Thr Leu Ser Arg 5

<210> 117

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 117

ser Ser Asp Trp Cys Phe Gly Trp Gly Gly Trp Cys Ala Ser Glu Ala 10

Val Ser Arg

<210> 118

<211> 19

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 118

Ser Arg Asn Trp Asp Trp Ala Ala Leu Glu Leu Leu Pro Tyr Pro His 10

Pro Ser Arg

<210> 119

<211> 15

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 119

Ser Ser Leu Thr Ser Arg Asp Phe Gly Ser Trp Tyr Ala Ser Arg

<210> 120

<211> 15

<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Arbitrary
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<400> 120
Ser Arg Ser Pro Ile Leu Thr His Leu Leu Ser Leu Gly Ser Arg
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<210> 121
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
    peptide
<400> 121
Ser Ser Thr Gly Ile Leu Trp Lys Leu Leu Thr Ala Glu Ser Arg
                  5
                                     10
<210> 122
<211> 15
<212> PRT
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     peptide
<400> 122
Ser Ser His Gly Ile Leu Trp Arg Leu Leu Ser Glu Gly Ser Arg
                                     10
<210> 123
<211> 15
<212> PRT
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<400> 123
Ser Arg Ser Asp Ser Ile Leu Trp Arg Met Leu Ser Glu Ser Arg
<210> 124
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<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 124
Ser Arg Leu Val Ala Leu Leu Lys Ser Pro Trp Ser Val Ser Arg
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 <400> 125
 Ser Arg Leu Glu Glu Leu Leu Met Asp Phe Trp Arg Ser Arg
                                       10
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<212> PRT
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<220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 126
 Ser Ser Lys Leu Trp Gln Leu Leu Ser Ser Pro Ile Asp Ser Arg
                                      10
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      peptide
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 Ser Ser Lys Leu Tyr Cys Leu Leu Asp Glu Ser Tyr Cys Ser Arg
                                        10
                    5
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  <211> 19
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  <223> Description of Artificial Sequence: Arbitrary
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peptide
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Val Ser Arg

<210> 129

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Arbitrary
peptide

<400> 129

Ser Ser Arg Leu Leu Ala Cys Glu Leu Met Tyr Glu Asp Ala Asp Val 1 5 10 15

Cys Ser Arg

<210> 130

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Arbitrary
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<400> 130

His Ser His Ser Pro Leu Leu Met Ala Leu Leu Ala Pro Pro Gly Gly

<210> 131

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Arbitrary
peptide

<400> 131

Ser Arg Leu Glu Tyr Tyr Leu Arg Leu Gly Thr Tyr Glu Ser Arg 1 5 10 15

<210> 132

<211> 15

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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
·<400> 132
Ser Ser Cys Leu Arg Glu Ile Leu Leu Tyr Gly Ala Cys Ser Arg
                                      10
<210> 133
<211> 19
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 133
Ser Ser Arg Thr Ala Glu Asp Tyr Cys Phe Phe Ala Asp Asp Tyr Trp
Cys Ser Arg
<210> 134
<211> 19
 <212> PRT
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      peptide
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                                                           15
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 Cys Ser Arg
 <210> 135
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 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 <400> 135
 Ser Ser Tyr Lys Pro His Ser Leu Leu Glu Trp His Leu Leu Gly Gly
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15

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Thr Ser Arg
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<210> 136
<211> 15
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<213> Artificial Sequence
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<400> 136
Ser Arg Leu His Cys Leu Leu Asp Ser Ser Tyr Cys Ser Ser Arg
<210> 137
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 137
Ser Arg Leu His Cys Leu Leu Asp Ser Ser Tyr Cys Ser Ser Arg
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<210> 138
<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 138
Ser Ser Trp Pro Asn Pro Thr Phe Trp Glu Arg Gln Leu Ser Arg
                   5
                                      10
 1 .
<210> 139
<211> 14
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
    · peptide
<400> 139
 Ser Tyr Ser Lys Glu Trp Phe Glu Glu Arg Leu Asn Ser Arg
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<210> 140
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 140
Ser Ser Ser Met Met Arg Glu Phe Phe Glu Arg Glu Leu Ser Arg
 <210> 141
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 141
 Ser Ser Gly Leu Pro Pro Asn Phe Glu Arg Met Leu Lys Ser Arg
                                       10
 <210> 142
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 142
 Ser Ser Gly Pro Trp Leu Met His Tyr Leu Gly Gly Gly Ser Arg
                                       10
 <210> 143
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 143
  Ser Ser Thr Ser Trp Leu His His Tyr Leu Met Gly Thr Ser Arg
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<210> 144
<211> 15
<212> PRT
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<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 144
Ser Arg Gly Gly Glu Cys Leu Gly Pro Trp Cys Leu Ser Arg
                                     10
<210> 145
<21.1> 19
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 145
Ser Ser Glu Ala Cys Val Gly Arg Trp Met Leu Cys Glu Gln Leu Gly
                                      10
Val Ser Arg
<210> 146
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 146
Ser Ser Gln Val Trp Pro Gly Pro Trp Arg Leu Val Glu Ser Arg
                                      10
<210> 147
<211> 15
<212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 <400> 147
 Ser Ser Ser Leu Gly Pro Trp Arg Leu Ser Glu Leu Glu Ser Arg
                   5
                                      10
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<210> 148

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<211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
<223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 148
  Ser Ser Ser Gly Pro Trp Arg Trp Gly Leu Ser Ile Glu Ser Arg
                                                            15
                   5 .
  <210> 149
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 149
  Ser Arg Glu Cys Val Gly Gly Trp Cys Leu Ala Glu Leu Ser Arg
  <210> 150
  <211> 15
<212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  Ser Ser Ile Pro Pro Arg Ser Trp Trp Leu Ser Gln Leu Ser Arg
                                        10
                                                             15
                   . 5
  <210> 151
   <211> 15
   <212> PRT
   <213> Artificial Sequence
   <223> Description of Artificial Sequence: Arbitrary
        peptide
   <400> 151
   Ser Ser Trp Pro Gly Ala Glu Trp Phe Lys Glu Gln Leu Ser Arg
   <210> 152
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<211> 15
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<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 152
Ser Ser Lys Leu Tyr Cys Leu Leu Asp Glu Ser Tyr Cys Ser Arg
                                      10
<210> 153
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 153
His Ser Tyr Ser Ser His Pro Leu Leu Leu Ser Tyr Leu Trp Gly Gly
                                                           15
                  5
                                       10
<210> 154
<211> 16
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 154
His Ser Trp Leu Gly Pro Trp Arg Leu Ser Ser Ile Asp Leu Gly Gly
                                       10
<210> 155
<211> 16
<212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 155
His Ser Thr Asp Met Gly Trp Leu Arg Pro Trp Arg Leu Leu Gly Gly
                                       10
 <210> 156
<211> 15
<212> PRT
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
<400> 156
Ser Ser Val Phe Thr Ile Met Asp Gly Lys Val Ala Leu Ser Arg
     5
                                      10
<210> 157
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Pro Tyr Cys Leu Gly Asp Val Trp Cys Leu Asp Ser Arg
<210> 158
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 158
Ser Arg Glu Trp Glu Asp Gly Phe Gly Gly Arg Trp Leu Ser Arg
                                     10
<210> 159
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 159
Ser Ser Trp Asn Ser Arg Glu Phe Phe Leu Ser Gln Leu Ser Arg
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<210> .160
<211> 15
<212> PRT
<213> Artificial Sequence
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<220>
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 160
 Ser Ser Thr Thr Met Phe Asp Phe Phe Tyr Glu Arg Leu Ser Arg
                                       10
 <210> 161
 <211> 15.
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 161
 Ser Ser Ala Arg Pro Trp Trp Leu Gln Phe Glu Gly Ser Ser Arg
                  5
 <210> 162
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 Ser Ser Gln Glu Glu Trp Leu Leu Pro Trp Arg Leu Ala Ser Arg
                   5
                                       10
 <210> 163
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <220>
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 163
 Ser Arg Leu Pro Pro Ser Val Phe Ser Met Cys Gly Ser Glu Val Cys
 Leu Ser Arg
  <210> 164
  <211> 19
  <212> PRT
  <213> Artificial Sequence
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<220>

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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 164
Ser Ser Gly Pro Phe Tyr Val Gly Gly Met Leu Trp Pro Ala Asp Cys
                  5
 1
                                      10
Leu Ser Arg
<210> 165
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide ·
<400> 165
Ser Arg Glu Gly Trp Met Gly Pro Trp Arg Leu Ala Asp Ser Arg
 1
                                      10
<210> 166
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 166
Ser Arg Asn Glu Cys Ile Gly Pro Trp Cys Leu Thr Ile Ser Arg
                                      10
<210> 167
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 167
Ser Ser Pro Gly Ser Arg Glu Trp Phe Lys Asp Met Leu Ser Arg
                                     10
1
<210> 168
<211> 15
<212> PRT
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<210> 172 <211> 19

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<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 168
Ser Ser Val Ala Ser Arg Glu Trp Trp Val Arg Glu Leu Ser Arg
                                      10
<210> 169
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 169
Ser Arg Met Phe Gln Val Cys Gly Asp Glu Val Cys Leu Arg Ser Arg
                                      10
<210> 170
<211> 16
<212> PRT
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     peptide
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<210> 171
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 171
Ser Arg Leu Asn Gly Val Phe Cys His Asp Ser Ser Asp Leu Trp Val
                                     10
                                                          15
Cys Ser Arg
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peptide

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<212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
        peptide
 <400> 172
 Ser Arg Pro Gly Cys Leu Arg Gly Val Trp Cys Leu Ala Asp Thr Pro
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                                       10
                                                            15
 Pro Ser Arg
 <210> 173
 <211> 19
 <212> PRT
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 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 173
. Ser Ser Arg Leu Val Pro His Ser Phe Trp Leu Asp Gly Leu Met His
                                       10
 Gly Ser Arg
 <210> 174
 <211> 19
 <212> PRT
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 <400> 174
 Ser Ser Ile Ser Thr Tyr His Met Gly Glu Trp Phe Tyr Ala Met Leu
                   5
                                       10
                                                           15
 Ser Ser Arg
 <210> 175
 <211> 18
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
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```
<223> Description of Artificial Sequence: Arbitrary
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<400> 187
Ser Ser Ile Lys Asp Phe Pro Asn Leu Ile Ser Leu Leu Ser Arg
<210> 188
<211> 19 '
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<220>
<223> Description of Artificial Sequence: Arbitrary.
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<400> 188
Ser Ser Gly Ser Ser Ala Gly Arg Leu Met Mer Leu Leu Gln Asp Gly
                 . 5
Val Ser Arg
<210> 189
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
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      peptide
<400> 189
Ser Arg Glu Gly Leu Leu Met Arg Leu Leu Ile Gly Asp Ser Arg
                                      10
<210> 190
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
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      peptide
<400> 190
Ser Ser His Cys His Thr Arg Leu Cys Ser Leu Leu Thr Ser Arg
                   5
                                                           15
<210>.191
 <211> 15
<212> PRT
 <213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 191
Ser Ser Arg Leu Leu Cys Leu Leu Asp Ala Gly Gln Cys Ser Arg
                  5
<210> 192
<211> 15
<212> PRT
<213> Artificial Sequence
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Ser Arg Asn Leu Leu Cys Leu Leu Asp Gln Glu Ala Cys Ser Arg
<210> 193
<211> 14
<212> PRT
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<220>
<223> Description of Artificial Sequence: Arbitrary
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<400> 193
Ser Ser Leu Lys Cys Leu Leu Asn Ser Asn Phe Cys Ser Arg
<210> 194
<211> 19
<212> PRT
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     peptide
<400> 194
Ser Ser Leu Lys Cys Leu Leu Gln Ser Ser Pro Gln Lys Gln Pro Phe
Cys Ser Arg
<210> 195
<211> 15
<212> PRT
<213> Artificial Sequence
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<220>

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<400> 195
Ser Ser Arg Thr Leu Leu Glu His Tyr Leu Leu Gly Gly Ser Arg
<210> 196
<211> 15
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<213> Artificial Sequence
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      peptide
<400> 196
Ser Ser Ala Gly Leu Leu Glu Asp Met Leu Arg Ser Arg Ser Arg
                                      10
<210> 197
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
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Ser Ser Arg Cys Ser Ser Leu Leu Cys Glu Met Leu Ile Gln Thr Lys
                  5
Glu Ser Arg
<210> 198
<211> 19
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 198
Ser Ser Leu Gln Ala Gly Ser Trp Leu Met His Tyr Leu Arg Gly Gly
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Asp Ser Arg
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<210> 199
 <211> 15
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 <223> Description of Artificial Sequence: Arbitrary
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 <400> 199
Ser Arg Pro Glu Gly Ser Ser Trp Leu Leu His Tyr Leu Ser Arg
                                                          15
<210> 200
<211> 15
<212> PRT
<213> Artificial Sequence
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     peptide
<400> 200
Ser Ser Arg Thr Leu Leu Glu His Tyr Leu Leu Gly Gly Ser Arg
                                      10 .
<210> 201
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
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     peptide
<400> 201
Ser Arg Trp Trp Leu Asp Asp His Glu Leu Leu Leu Tyr Ser Ser Arg
                5
                                      10
<210> 202
<211> 19
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 202
Ser Ser Arg Thr Leu Tyr Cys His Leu Thr Ser Ser Asn Pro Glu Trp
                  5
                                    10
Cys Ser Arg
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<210> 203
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 <212> PRT
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<220>
 <223> Description of Artificial Sequence: Arbitrary
<400> 203
Ser Ser Thr Arg Leu Met Cys Trp Leu Gly Ser Ala Asp Thr Ser His
Cys Ser Arg
<210> 204
<211> 19
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 204
Ser Ser Tyr Asp Trp Gln Cys Pro Ser Trp Tyr Cys Pro Ala Pro Pro
Ser Ser Arg
<210> 205
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 205
Ser Ser Thr Thr Trp Arg Cys Pro Glu Trp Tyr Cys Gly Ser Arg
                                      10
<210> 206
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
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<400> 206
Ser Ser Trp Asp Phe Arg Val Pro Trp Trp Tyr Asn Asn Ser Arg
                                       10
<210> 207
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 207
Ser Ser Gln Trp Gln Ala Pro Trp Trp Tyr Ile Asp Ala Ser Arg
<210> 208
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Ser Arg Pro Ser Phe Thr Ile Pro Trp Trp Phe Asp Asp Pro Ser
                  5
                                      10
Arg Ser Arg
<210> 209
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
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Ser Ser Tyr Glu Ile Pro Lys Trp Ala Leu Gln Trp Leu Ser Arg
                 5
<210> 210
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     paptida
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Ser Ser Leu Asp Leu Ser Gln Phe Pro Met Thr Ala Ser Phe Leu Arg
                  5
Glu Ser Arg
<210> 211
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 211
 Ser Ser Asn His Gln Ser Ser Arg Leu Ile Glu Leu Leu Ser Arg
 <210> 212
 <211> 14
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 212
 Ser Ala Pro Arg Ala Thr Ile Ser His Tyr Leu Met Gly Gly
                    5
   1 . .
  <210> 213
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 213
  Ser Ser Trp Asp Met His Gln Phe Phe Trp Glu Gly Val Ser Arg
                                   . 10
  <210> 214
  <211> 19
  <212> . PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
```

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peptide
```

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Ser Arg Leu Pro Pro Ser Val Phe Ser Met Cys Gly Ser Glu Val Cys
<400> 214
                  5
```

Leu Ser Arg

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<210> 215 <211> 15 <212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

Ser Ser Pro Gly Ser Arg Glu Trp Phe Lys Asp Met Leu Ser Arg 10 5

<210> 216 <211> 15

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 216 Ser Ser Glu Tyr Cys Phe Tyr Trp Asp Ser Ala His Cys Ser Arg 10 1

<210> 217

<211> 15

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 217

Ser Ser Leu Thr Ser Arg Asp Phe Gly Ser Trp Tyr Ala Ser Arg 10 5

<210> 218

<211>. 15

<212> PRT

<213> Artificial Sequence

<220>

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<223> Description of Artificial Sequence: Arbitrary
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Ser Arg Thr Trp Glu Ser Pro Leu Gly Thr Trp Glu Trp Ser Arg
                  5
<210> 219
<211> 15
<212> PRT
<213> Arcificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
Ser Arg Glu Trp Glu Asp Gly Phe Gly Gly Arg Trp Leu Ser Arg
 <210> 220
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 220
 Ser Ser Leu Asp Leu Ser Gln Phe Pro Met Thr Ala Ser Phe Leu Arg
 Glu Ser Arg
 <210> 221
 <211> 19
 <212> PRT
 <213> Artificial Sequence
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       peptide
  Ser Ser Glu Ala Cys Val Gly Arg Trp Met Leu Cys Glu Gln Leu Gly
  <400> 221
                                        10
   1
  Val Ser Arg
  <210> 222
<211> 15
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```
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Ala Gly Leu Leu Ser Asp Leu Leu Glu Gly Lys Ser Arg
<400> 222
                  5
 <210> 223
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 Ser Ser Arg Ser Leu Leu Arg Asp Leu Leu Met Val Asp Ser Arg
 <400> 223
                   5
 <210> 224
 <211> 15
 <212> PRT
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       peptide
  Ser Ser Asn Lys Leu Leu Tyr Asn Leu Leu Lys Met Glu Ser Arg
  <400> 224
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   1
  <210> 225
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  Ser Ser Lys Ser Leu Leu Leu Asn Leu Leu Ser Thr Pro Ser Arg
                                        10
                     5
   <210> 226
   <211> 16
   <212> PRT
   <213> Artilicial Sequence
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<220>

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<223> Description of Artificial Sequence: Arbitrary
     peptide
His Ser Phe Pro Arg Glu Ser Leu Leu Val Arg Leu Leu Gln Gly Gly
<210> 227
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Leu Glu Met Leu Leu Arg Ser Glu Thr Asp Phe Ser Arg
 1
 <210> 228
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 Ser Arg Leu Glu Glu Leu Leu Lys Trp Gly Ser Val Thr Ser Arg
 <400> 228
                                      10
 1
 <210> 229
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 229
  Ser Arg Leu Glu Gln Leu Leu Lys Glu Glu Phe Ser Tyr Ser Arg
                                       10
                    5
  <210> 230
  <211> .15
  <212> PRT
  <213> Artificial Sequence
  くて202
```

```
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 230
Ser Arg Leu Glu Gln Leu Leu Arg Ser Glu Pro Asp Phe Ser Arg
                                     10
                 5
<210> 231
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 231
Ser Arg Leu Glu Asp Leu Leu Arg Ala Pro Phe Thr Thr Ser Arg
                                      10
                   5
<210> 232
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 232
 Ser Arg Leu Glu Ser Leu Leu Arg Phe Gly Gln Leu Asp Ser Arg
                                                           15
                                       10
                   5
 <210> 233
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 233
 Ser Ser Arg Leu Leu Ser Leu Leu Val Gly Asp Phe Asn Ser Arg
                                       10
                   · 5
  <210> 234
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
        Poptide
```

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From-BROWDY NEIMARK

```
Ser Arg Leu Glu Glu Leu Leu Gly Thr Asn Arg Asp Ser Arg
                  5
<210> 235
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
Ser Arg Leu Glu Glu Leu Leu Met Asp Phe Trp Arg Ser Arg
<400> 235
                                     10
                  5
 <210> 236
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 Ser Arg Leu Lys Glu Leu Leu Leu Pro Thr Asp Leu Ser Arg
                                      10
              , 5
 <210> 237
 <211> 15
 <212> PRT
 <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 237
  Ser Arg Leu Glu Cys Leu Leu Glu Gly Arg Leu Asn Cys Ser Arg
                                      10
                   5
  <210> 238
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 238
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Ser Ser Lys Leu Tyr Cys Leu Leu Asp Glu Ser Tyr Cys Ser Arg
                                    10
<210> 239
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 239
Ser Arg Leu Ser Cys Leu Leu Met Gly Phe Glu Asp Cys Ser Arg
                  5
<210> 240
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 240
 Ser Ser Lys Leu Ile Arg Leu Leu Thr Ser Asp Glu Glu Leu Ser Arg
 <210> 241
 <211> 16
 <212> PRT
 <213> Artificial Sequence
 <220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
 <400> 241
 Ser Ser Arg Leu Met Glu Leu Leu Gln Glu Gly Gln Gly Trp Ser Arg
                                      10
                  5
 <210> 242
 <211> 15
<212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      .peptide
 Ser Ser Asn His Gln Ser Ser Arg Leu Ile Glu Leu Leu Ser Arg
                                       10
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<210> 243

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<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
Ser Ser Arg Leu Trp Gln Leu Leu Ala Ser Thr Asp Thr Ser Arg
                                      10
                  5
<210> 244
<211> 15
<212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide .
 Ser Ser Lys Leu Trp Gln Leu Leu Ser Ser Pro Ile Asp Ser Arg
 <210> 245
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 Ser Arg Leu Val Ala Leu Leu Lys Ser Pro Trp Ser Val Ser Arg
  <400> 245
                                       10
  <210> 246
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  Ser Ser Asn Ser Met Leu Trp Lys Leu Leu Ala Ala Pro Ser Arg
                                        10
                     5
    ı
```

```
<210> 247
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 247
Ser Ser Lys Thr Leu Trp Arg Leu Leu Glu Gly Glu Arg Ser Arg
                                     10
                  5
<210> 248
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 248
Ser Arg Ala Gly Pro Val Leu Trp Gly Leu Leu Ser Glu Ser Arg
<210> 249
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 249
Ser Arg Ser Pro Ile Leu Thr His Leu Leu Ser Leu Gly Ser Arg
                   5
<210> 250
<211> 15
 <212> PRT '
 <213> Artificial Sequence
<220>
 <223> Description of Artificial Sequence: Arbitrary
     peptide
 <400> 250
 Ser Ser Thr Gly Ile Leu Trp Lys Leu Leu Thr Ala Glu Ser Arg
 <210> 251
 <211> 15
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From-BROWDY NEIMARK.

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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 251
Ser Ser His Gly Ile Leu Trp Arg Leu Leu Ser Glu Gly Ser Arg
<210> 252
<211> 11
<212> PRT
<213> Human steroid receptor coactivator la
<400> 252
Lys Leu Val Gln Leu Leu Thr Thr Thr Ala Glu
                  5
  1
<210> 253
<211> 11
<212> PRT
<213> Human steroid receptor coactivator la
              · . .
 <400> 253
 Ile Leu His Arg Leu Leu Gln Glu Gly Ser Pro
                 5
 <210> 254
 <211> 11
 <212> PRT
 <213> Human steroid receptor coactivator la
 <400> 254
 Leu Leu Arg Tyr Leu Leu Asp Lys Asp Glu Lys
                  5
 <210> 255
 <211> 8.
 <212> PRT
 <213> Human steroid receptor coactivator la
 <400> 255
 Leu Leu Gln Gln Leu Leu Thr Glu
 <210> 256
 <211>,11
  <212> PRT
 <213> Mouse cAMP-responsive element (CREB)-binding protein
 <400> 256
```

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Gln Leu Ser Glu Leu Leu Arg Gly Gly Ser Gly
                5
<210> 257
<211> 11
<212> PRT
<213> Mouse cAMP-responsive element (CREB)-binding protein
<400> 257
Gln Leu Val Leu Leu His Ala His Lys Cys
<210> 258
<211> 11
<213> Mouse cAMP-responsive element (CREB)-binding protein
<212> PRT
<400> 258
Tyr Leu Glu Gly Leu Leu Met His Gln Ala Ala
                  5
  1
 <210> 259
 <211> 11
 <212> PRT
 <213> Mouse cAMP-responsive element (CREB)-binding protein
 Leu Leu Ala Ser Leu Leu Gln Ser Glu Ser Ser
                   5
 <210> 260
 <211> 11
 <212> PRT
 <213> Mouse cAMP-responsive element (CREB)-binding protein
 <400> 260
 His Leu Lys Thr Leu Leu Lys Lys Ser Lys Val
                                       10
 <210> 261
 <211> 11
  <212> PRT
  <213> Human RIP140
  <400> 261
  Gln Leu Ala Leu Leu Leu Ser Ser Glu Ala His
                    5
  1
  <210> 262
  <211> 11
  <212> PRT
<Z13> Human RIP140
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<400> 262
Leu Leu Leu His Leu Leu Lys Ser Gln Thr Ile
               5
<210> 263
<211> 11
<212> PRT
<213> Human RIP140
<400> 263
Leu Leu Gln Leu Leu Gly His Lys Asn Glu
1 5 10
<210> 264
<211> 11
<212> PRT
<213> Human RIP140
 <400> 264
Val Leu Gln Leu Leu Ely Asn Pro Lys Gly
               5 .
                                 10
 1
 <210> 265
 <211>:11
 <212> PRT
 <213> Human RIP140
 <400> 265
 Leu Leu Ser Arg Leu Leu Arg Gln Asn Gln Asp
             . 5
 1
 <210> 266
 <211> 11
 <212> PRT
 <213> Human RIP140
<400> 266
Val Leu Lys Gln Leu Leu Ser Glu Asn Cys
  1 5
 <210> 267
 <211> 14
 <212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      .peptide
  <400> 267
  Ser Ser Asn His Gln Ser Arg Leu Ile Glu Leu Leu Ser Arg
```

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<210> 268
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
His Val Tyr Gln His Pro Leu Leu Ser Leu Leu Ser Ser Glu His
                                      10
                   5
 Glu Ser Gly
 <210> 269
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
      pepride
 <400> 269
 His Val Glu Met His Pro Leu Leu Met Gly Leu Leu Met Glu Ser Gln
                                     10
                  5
 Trp Gly Ala
 <210> 270
  <211> 19
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Arbitrary
        peptide
  <400> 270
  Gln Glu Ala His Gly Pro Leu Leu Trp Asn Leu Leu Ser Arg Ser Asp
    1
  Thr Asp Trp
  <210>.271
  <211> 19
  <212> PRT
  <213> Artificial Sequence
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<400> 274

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```

```
<223> Description of Artificial Sequence: Arbitrary
     peptide
Gly His Glu Pro Leu Thr Leu Leu Glu Arg Leu Leu Met Asp Asp Lys
                                    10
Gln Ala Val
<210> 272
<211> 19
<212> PRT
<213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 Leu Pro Tyr Glu Gly Ser Leu Leu Leu Lys Leu Leu Arg Ala Pro Val
                                      10
  1
 Glu Glu Val
 <210> 273
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 273
 Ser Gly Trp Glu Asn Ser Ile Leu Tyr Ser Leu Leu Ser Asp Arg Val
                              . 10
                   5
  Ser Leu Asp
  <210> 274 .
  <211> 19
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
        peptide
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Ala His Gly Glu Ser Ser Leu Leu Ala Trp Leu Leu Ser Gly Glu Tyr

```
Ser Ser Ala
```

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<210> 275
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
Gly Val Phe Cys Asp Ser Ile Leu Cys Gln Leu Leu Ala His Asp Asn
<400> 275
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Ala Arg Leu

<210> 276 <211> 19 <212> PRT <213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 276 His His Asn Gly His Ser Ile Leu Tyr Gly Leu Leu Ala Gly Ser Asp 10

10

Ala Pro Ser

<210> 277 <211>.19 <212> PRT <213> Artificial Sequence

<223> Description of Artificial Sequence: Arbitrary peptide

<400> 277 Leu Gly Glu Arg Ala Ser Leu Leu Asp Met Leu Leu Arg Gln Glu Asn 15

Pro Ala Trp

<210> 278 <211> 19

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<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 278
Ser Gly Trp Asn Glu Ser Thr Leu Tyr Arg Leu Leu Gln Ala Asp Ala
                                      10
Phe Asp Val
<210> 279
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence; Arbitrary
      peptide
 <400> 279
 Pro Ser Gly Gly Ser Ser Val Leu Glu Tyr Leu Leu Thr His Asp Thr
                                      10
 Ser Ile Leu
 <210> 280
 <211> 19
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 280
 Gly Ser Glu Pro Lys Ser Arg Leu Leu Glu Leu Ser Ala Pro Val
                                      10
 Thr Asp Val
 <210> 281
 <211> 19
 <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
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<210> 282 <211> 19 <212> PRT <213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Arbitrary peptide

<400> 282 Val Glu Ser Gly Ser Ser Arg Leu Met Gln Leu Leu Met Ala Asn Asp 10 15

Leu Leu Thr

<210> 283
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence; Arbitrary

Ala Val Trp

peptide

<210> 284 <211> 19. <212> PRT <213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Arbitrary
 peptide

Glu Arg Glu

.5

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<210> 285
<211> 12
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
Met Ser Trp Tyr Glu Phe Met Thr Glu Glu Ser Met
<400> 285
                                    10
                  5
<210> 286
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Arbitrary
      peptide
 Ala Lys His Asp Leu Ser Trp Tyr Glu Phe Leu Gln Leu Pro Ile
                                     10
 <210> 287
 <211> 17
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
  Ser Arg Leu Ser Trp Trp Glu Phe Leu Gly Ala Ser Asp Cys Gly Thr
 <400> 287
                          10 15
                  5
  Cys
  <210> 288
  <211> 11
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: Arbitrary
       peptide
  <400> 288
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<210> 309
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From-BROWDY NEIMARK

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<211> 17
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                   5
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### IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:	) ART UNIT: 1627
PAIGE et al	) Examiner: T. Wessendorf
Appln. No.: 09/429,331	) Washington, D.C.
Filed: October 28, 1999	) February 27, 2001
For: METHOD OF PREDICTING THE ABILITY OF COMPOUNDS TO MODULATE THE BIOLOGICAL ACTIVITY OF RECEPTORS	Atty.Docket: PAIGE=1D ) ) )

SUPPLEMENTAL RESPONSE TO "SEQUENCE LISTING" REQUIREMENT

Honorable Commissioner of Patents Washington, D.C. 20231

Sir:

Supplementing the response filed February 20, 2001, please further amend the application as follows:

### IN THE SPECIFICATION

At page 162, line 34, replace "K = C or T" with --K = G or T--.

Please replace the present page 237 with the enclosed substitute page, which provides SEQ ID NOs:316-360 for the sequences on this page. If there was no page numbered "237" in the application as filed, please insert this page between pages "236" and "238".

The enclosed "Sequence Listing" pages 1, 4 and 79-90 replace pages 1, 4 and 79 of the "Sequence Listing" submitted February 20, 2001.

In re Appln. No -09/429, 331

#### REMARKS

1. At the time the February 20, 2001, response was prepared, counsel's file copy of the specification was missing page 237. Hence, the sequences appearing on that page were not incorporated into the Sequence Listing filed on that date.

Since counsel received a postcard receipt (copy enclosed) acknowledging the filing of a 293 page specification, counsel assumes that page 237 was missing only from counsel's file copy and not from the original filed with the PTO.

If counsel is mistaken, inserting page 237 at this time does not constitute the addition of "new matter". At page 1, lines 3-10, it is stated:

This application is a continuation-in-part of PCT/US99/06664, filed March 26, 1999, which is a continuation-in-part of 60/115,345, filed January 8, 1999, which is a continuation-in-part of Paige et al., Serial No. 60/099,656, filed September 9, 1998, which is a continuation-in-part of Paige et al., Serial No. 60/082,756, filed April 23, 1998. All of the above applications are hereby incorporated-by-reference.

Page 237 of this application sets forth Table 1, and part of Table 2. It is identical to page 152 of the above-identified, incorporated-by-reference PCT application. Hence, even if inadvertently omitted from this application as filed, it can be provided without adding "new matter".

In re Appln. No 39/429,331

- 2. At page 162, we correct an obvious typographical error in the identification of ambiguous nucleotide "K", which denotes "G" or "T", not "C" or "T". See MPEP \$2422, page 2400-20, Table 1. The NNK codon, specified at page 162, line 33, encodes all 20 amino acids. If the third position were C/T (Y), instead of G/T (K), then Met (ATG), Trp (TGG), Ser (TCA, TCG), Gln (CAA, CAG), Lys (AAA, AAG) and Gly (GAA, GAG) would not be encoded, inconsistent with the identification of X in LXXLL (page 162, line 29) as "any AA". This error was also corrected on page 4 of the Sequence Listing at <223> in SEQ ID NO:14.
  - 3. Applicants hereby submit the following:
  - [XX] an amendment to the paper copy of the "Sequence Listing" submitted on February 20, 2001, the amendment being in the form of substitute pages 1 and 79 and new pages 80-90;
  - [XX] the Sequence Listing in computer readable form, complying with \$1.821(e) and \$1.824, including, if an amendment to the paper copy is submitted, all previously submitted data with the amendment incorporated therein;
- [XX] 4. The description has been amended to comply with \$1.821(d).

5. The undersigned attorney or agent hereby states as follows:

- (a) this submission is not believed to include new matter [\$1.821(g)];
- (b) the contents of the paper copy (as amended, if applicable) and the computer readable form of the Sequence Listing, are believed to be the same [\$1.821(f) and \$1.825(b)];
- (c) if the paper copy has been amended, the amendment is believed to be supported by the specification and is not believed to include new matter [\$1.825(a)]; and

Respectfully submitted,

BROWDY AND NEIMARK

Attorneys for Applicant(s)

By:

Iver P. Cooper

Registration No. 28,005

IPC:al
624 Ninth Street, N.W.
Washington, D.C. 20001
Telephone No.: (202) 628-5197

Facsimile No.: (202) 737-3528 F:\,N\Nova\PaigelD\Pto\SequenceResponse.doc

### Enclosures:

Paper Sequence Listing pp. 1,4 and 79-90 Substitute CRF Substitute page 237 Page 152 of PCT/US99/06664 Copy of stamped postcard receipts

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PCT/US99/06664

### Table 1

Peptides the Bind to the Unliganded (unactivated)

	Estrogen	Receptor	Sequence	Phage #
5	s R		TWEWSR .	4 48
	S A S S	WVRLSD	FPWGVSR	1 2
	S S S S			3
10	\$ S	WVLLRI		31 29
	s s	CKWYE	CSGLWSR	7 35
	S S S R	NLCFFV	N D D E Y C S R	41 47
15.	н н	IHRHPAI	ирнтусс	~ /

# Table 2

# Peptides that Bind to the Estradiol Activated

	Receptor	•
	Sequence	Phage #
		1/2
20	SKAGILISTANDOD	б
	SSRUGA	22
		23
	2000	42
		3
25		11
		21
-	2 / 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	27
•	2 K B B Z B B B B B B B	28
		29
30	SRLESLLRFGQLDSR	19/20
	SSRLLSLLVGDFNSR	30
	SRLEELLLGTNRDSR	. 15
	SRLKELLLPTDLSR	34
	SRLECLLEGRLNCSR	35
35	SSKLYCLLDESYC SR	36
	SRLSCLLMGFEDCSR	37
	SSKLIRLLTSDEELSR	
	SSRLMELLQEGQGWSR	40
	SSNHQSSRLIELLSR	4
40	SSRLWQLLASTDTSR	16
-10	SSNSMLWKLLAAPSR	13/14
•	SSKTLWRLLEGERSR	17
	SRAGPVLWGLLSESE	32 .
	SSLTSRDFGSWYASR	5
4 =	SSWVRLSDFPWGVSR	24/25
45	SSEYCFYDSAHCSR	33 .
	SRSLLECHLMGNCSR	7
	SSELLRWHLTRDTSR	8
		. 12
	B K B B a s a s a s a s a s a s a s a s a s	· 31
50		38/39
	33 2 3 1 2 1 2 1 2 1 2 1 2 1 2 1 2 1 2 1	41
	H S H G L M L L M M M M M M M M M M M M M M M	26
	SSAGGGAPAGSTPSK	

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Table 1

Peptides the Bind to the Unliganded (unactivated) Estrogen Receptor

		_	Sequence	sea id no:	Phage #
_	a n	0	TWEWSR	316	4
5	SR	APRTIS	HYLMGG	317	48
			FPWGVSR	318	ı
	SS		FPWGVSR	319	2
	8 8	W D W = -		320	3
	SS	M # 17 = 20		321	31
10	S S	M ^ T = 22	LPWGSR	322	29
•	S S	WVJ	CSGLWSR	323	7
	\$ \$			324	、35
	S S	GICFFW		325	41
	SR	NLCFFW		326	47
1.5	. нн	HRHPAH	PHTYGG	<b>.</b>	

## Table 2

# Peptides that Bind to the Estradiol Activated Receptor

	-	SEB IDNO:	Phage #
	Sequence		
20	SRAGLLSDLLEGKSR	327	1/2
20		328	6
	S S R S L L R D L L M V D S R S S N K L L Y N L L K M E S R	329	22
	SSKSLLLNLLSTPSR	330	23
*		331	42
	H 2 E E K I D I I I I I I I I I I I I I I I I I	332	3
25	SKIII	333	11
	SKHALLINGTOVOR	334	21
		335	. 27
	SKUDQUIII	336	28
	SKHEDHAAAAAA	337	29
30		338	19/20
	S C T T D C P	339	30
	SKUMBATTTTTTTTTTT	340	15
	SKIKI		34
	SKUBCUHA	341 342	35
35	227 1 1 1 1 1 2 2 2		36
	SRLSCLLMGFEDCSR	343	37
	SSKLIRLLTSDEELSR	344	40
	SSRLMELLQEGQGWSR	345	4
	SSNHQSSRLIELLSR	346	16
40	SSRLWOLLASTDTSR	347	
40	SSNSMLWKLLAAPSR	348	13/14
	SSKTLWRLLEGERSR	उपद	17
	CPAGPVLWGLLSESR	350	32
	CST.TSRDFGSWYASR	3 <b>5</b> 1	5
	SSWVRLSDFPWGVSR	352	24/25
45		353	33
		354	7
		355	8
		35b	12
			31
50	SKSDSLESS	357	38/39
	5529 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	358	41
	HSHGPLTLNLLRSSGG	359	26
	SSAGGGAPAGSTPSR	3 Lo	20

### SEQUENCE LISTING

- <110> PAIGE, Lisa A. MCDONNELL, Donald P. CHANG, Ching Yu NORRIS, John HAMILTON, Paul T. FOWLKES, Dana M. BARNETT, Tom , CHRISTIANSEN, Dale J. BUEHRER, Benjamin
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- <170> PatentIn Ver. 2.0
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- Gly Ser Gly Lys
- <210> 2
- <211> 8
- <212> PRT
- <213> Artificial Sequence

<400> 11

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                                      10
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      peptide
Ser Ser Pro Gly Ser Arg Glu Trp Phe Lys Asp Met Leu Ser Arg
                                      10
<210> 13
<211> 15
<212> PRT
<213> Artificial Sequence
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<400> 13
Ser Ser Leu Thr Ser Arg Asp Phe Gly Ser Trp Tyr Ala Ser Arg
<210> 14
<211> 88
<212> DNA
<213> Artificial Sequence
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      peptide library of Ex. 101.1
<220>
<223> N at each occurrence is A, C, G or T; K at each
      occurrence is G or T
<400> 14
agtgtgtgcc tcgagannkn nknnknnknn knnknnkctg nnknnkctgc tgnnknnknn 60
knnknnknnk nnktotagac tgtgcagt
<210> 15
<211> 15
<212> DNA
<213> Artificial Sequence
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<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 313
Ser Ser His Trp Ser Ser Asp Ser Ile Phe Pro Gly Phe Trp Tyr Ser
                                      10
Gly
<210> 314
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     peptide
<400> 314
Ser Arg Gly Gly Val Asp Leu Asp Ile Gly Asn Ser Ala
<210> 315
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       pepride
 <400> 315
 Glu Gly Glu Asp Val Arg Thr Arg Ile Ala Asn
   1
 <210> 316
 <211> 14
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       peptide
 <400> 316
 Ser Arg Trp Glu Ser Pro Leu Gly Thr Trp Glu Trp Ser Arg
                                       10
                   5
 <210> 317
 <211> 14
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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 317
Ser Ala Ala Pro Arg Thr Ile Ser His Tyr Leu Met Gly Gly
                                      10
<210> 318
<211> 15
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      peptide
<400> 318
Ser Ser Trp Val Arg Leu Ser Asp Phe Pro Trp Gly Val Ser Arg
                                                           15
<210> 319
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      peptide
<400> 319
Ser Ser Trp Asp Arg Leu Ser Asp Phe Pro Trp Gly Val Ser Arg
                   5
<210> 320
<211> 15
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      peptide
 <400> 320
 Ser Ser Trp Ile Arg Leu Arg Asp Leu Pro Trp Gly Glu Ser Arg
                   5
 <210> 321
 <211> 14
 <212> PRT
 <213> Artificial Sequence
```

<220>

```
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     peptide
<400> 321
Ser Ser Trp Val Leu Leu Arg Asp Leu Pro Trp Gly Ser Arg
                                     10
                  5
<210> 322 ·
<211> 14
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
   peptide
<400> 322
Ser Ser Trp Val Val Leu Arg Asp Leu Pro Trp Gly Ser Arg
                 5
<210> 323
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Arbitrary
     peptide
 <400> 323
Ser Ser Cys Lys Trp Tyr Glu Lys Cys Ser Gly Leu Trp Ser Arg
 <210> 324
 <211> 15
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Arbitrary
       peptide
 <400> 324
 Ser Ser Gly Ile Cys Phe Phe Trp Asp Gly Cys Phe Glu Ser Arg
 <210> 325
 <211> 15.
 <212> PRT
 <213> Artificial Sequence
 <220>
```

peptide

```
<223> Description of Artificial Sequence: Arbitrary
     peptide
<400> 325
Ser Arg Asn Leu Cys Phe Phe Trp Asp Asp Glu Tyr Cys Ser Arg
                                     10
<210> 326
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Arbitrary
    · peptide
<400>.326
His His Arg His Pro Ala His Pro His Thr Tyr Gly Gly
<210> 327
<211> 15
<212> PRT
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<220>
<223> Description of Artificial Sequence: Arbitrary
      peptide
<400> 327
Ser Arg Ala Gly Leu Leu Ser Asp Leu Leu Glu Gly Lys Ser Arg
                                      10
                  5
 <210> 328
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 <223> Description of Artificial Sequence: Arbitrary
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 Ser Ser Asn Lys Leu Leu Tyr Asn Leu Leu Lys Met Glu Ser Arg
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 Ser Ser Lys Ser Leu Leu Asn Leu Leu Ser Thr Pro Ser Arg
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Ser Arg Leu Glu Glu Leu Leu Gly Thr Asn Arg Asp Ser Arg
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Ser Ser Lys Leu Tyr Cys Leu Leu Asp Glu Ser Tyr Cys Ser Arg
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Ser Ser Arg Leu Trp Gln Leu Leu Ala Ser Thr Asp Thr Ser Arg
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Ser Ser Asn Ser Met Leu Trp Lys Leu Leu Ala Ala Pro Ser Arg
                                      10
                  5
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                                      10
<210> 350
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      peptide
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Ser Arg Ala Gly Pro Val Leu Trp Gly Leu Leu Ser Glu Ser Arg
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Ser Ser Glu Tyr Cys Phe Tyr Asp Ser Ala His Cys Ser Arg
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Ser Arg Ser Leu Leu Glu Cys His Leu Met Gly Asn Cys Ser Arg
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Ser Ser Glu Leu Leu Arg Trp His Leu Thr Arg Asp Thr Ser Arg
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Ser Arg Leu Glu Tyr Trp Leu Lys Trp Glu Pro Gly Pro Ser Arg
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Ser Arg Ser Asp Ser Ile Leu Trp Arg Met Leu Ser Glu Ser Arg
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Ser Ser Lys Gly Val Leu Trp Arg Met Leu Ala Glu Pro Val Ser Arg
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His Ser His Gly Pro Leu Thr Leu Asn Leu Leu Arg Ser Ser Gly Gly
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<400> 360
Ser Ser Ala Gly Gly Gly Ala Pro Ala Gly Ser Thr Pro Ser Arg
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